F Errors Corrected by the STIC Septems Branch 16	33
lumber: 09/336,3730 ENTEDED : 11	=11
Charged a mo nonverse	74
Changed the margins in cases where the sequence text was "wrapped" down to the next line.	
Edited a format error in the Current Application Data section, specifically:	
Edited the Current Application Data section with the actual current number. The number inputted applicant was  the prior application data; or other	by the
Added the mandatory heading and subheadings for "Current Application Data".	
Edited the "Number of Sequences" (ield. The applicant spelled out a number instead of using an i	nteger.
Changed the spelling of a mandatory field (the headings or subheadings), specifically:	· -
Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were	·e:
Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:	
Corrected subheading placement. All responses must be on the same line as each subheading. If applicant placed a response below the subheading, this was moved to its appropriate place.	the
Inserted colons after headings/subheadings. Headings edited included:	
Deleted extra, invalid, headings used by an applicant, specifically:	
Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at ☐ page numbers throughout text; ☐ other invalid text, such as	end of file;
Inserted mandatory headings, specifically:	<del></del>
Corrected an obvious error in the response, specifically:	
Edited identifiers where upper case is used but lower case is required, or vice versa.	
Corrected an error in the Number of Sequences field, specifically:	
A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	
Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field according due to a Patentin bug). Sequences corrected:	
Other: Inserted "hard returns" where needed this the sequence listing.	ughout
the sequence listing.	

<sup>\*</sup>Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

1633

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/356,575D

DATE: 11/17/2000 TIME: 09:26:43

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\11172000\I356575D.raw

```
2 <110> APPLICANT: Fall.x, Frits
        Hoeben, Robert
         Bout, Abraham
         Valerio, Domenico
         "an der Eb, Alex
         Schouten, Govert
9 <120> TITLE OF INVENTION: PACKAGING SYSTEMS
11 <130> FILE REFERENCE: 2578-3935US
13 <140> CURRENT APPLICATION NUMBER: US/09/356,575D
14 <141> CURRENT FILING DATE: 1999-07-19
15 <150> PRIOR APPLICATION NUMBER: US 08/793,170
17 <151> PRIOR FILING DATE: 1997-03-25
18 <150> PRIOR APPLICATION NUMBER: PCT/NL96/00244
19 <151> PRIOR FILING DATE: 1996-06-14
20 <150> PRIOR APPLICATION NUMBER: EP 95201611.1
21 <151> PRIOR FILING DATE: 1995-06-15
22 <150> PRIOR APPLICATION NUMBER: EP 95201728.3
23 <151> PRIOR FILING DATE: 1995-06-26
25 <160> NUMBER OF SEQ ID NOS: 22
27 <170> SOFTWARE: PatentIn version 3.0
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 21
31 <212> TYPE: DNA
32 <213> ORGANISM: Unknown
34 <220> FEATURE:
35 <223> OTHER INFORMATION: Derived from Adenovirus
 37 <400> SEQUENCE: 1
                                                                          21
38 cgtgtagtgt atttataccc g
41 <210> SEO ID NO: 2
42 <211> LENGTH: 21
 43 <212> TYPE: DNA
 44 <213> ORGANISM: Unknown
 46 <220> FEATURE:
 47 <223> OTHER INFORMATION: Derived from Adenovirus
 49 <400> SEQUENCE: 2
                                                                           2.1
 50 tegleactgg gtggaaagee a
 53 <210> SEQ ID NO: 3
 54 <211> LENGTH: 21
 55 <212> TYPE: DNA
 56 <213> ORGANISM: Unknown
 58 <220> FEATURE:
 59 <223> OTHER INFORMATION: Derived from Adenovirus
 61 <400> SEQUENCE: 3
                                                                           21
 62 tacccgccgt cctaaaatgg c
 65 <210> SEQ ID NO: 4
 66 <211> LENGTH: 20
 67 <212> TYPE: DNA
```

RAW SEQUENCE LISTING DATE: 11/17/2000 TIME: 09:26:43

-	
<pre>t8 &lt;213&gt; ORGANISM: Unkhown 70 &lt;220&gt; FLATURE: 71 &lt;223&gt; OTHER INFORMATION: Derived from Adenovirus</pre>	
73 <400> SEOJENCE: 4	20
7.1 raguettqay etqtaaaeqe	
77 <210> SLO 1D NO: 5	
78 <211> LENGIH: 21	
79 <212> TYPE: DNA	
80 <213> ORGANISM: Unknown	
82 <220> FEATURE: 83 <223> OFHER INFORMATION: Derived from Adenovirus	
83 (223) OTHER INFORMATION 55 85 (400) SEQUENCE: 5	21
86 decreated addressed t	
89 210> SEQ ID NO: 6	
90 \211> LENGTH: 20	
01 .212> TYPE: DNA	
92 <213> ORGANISM: Unknown	
45 -223> OTHER INFORMATION: Delived 110M. House	
97 .400> SEQUENCE: 9	20
98 gettqaqeee qagacatgte	
101 <210> SEQ TD NO: 7	
102 <211> LENGTH: 24	
103 <212> TYPE: DNA 104 <213> ORGANISM: Unknown	
106 <220> FEATURE: 107 <223> OTHER INFORMATION: Derived from Adenovirus	
109 <400> SEQUENCE: 7	24
110 cccctcgage tcaatctgta tett	
113 <21U> SEQ ID NO: 8	
114 <21.1> LENGTH: 27	
115 <212> TYPE: DNA	
116 <213> ORGANISM: Unknown	
118 <220> FEATURE: 119 <223> OTHER INFORMATION: Derived from Adenovirus	
119 <223> OTHER INFORMATION. BUTTLES	27
121 <400> SEQUENCE: 8 122 gggggatccg aacttgttta ttgcagc	2,
122 gggggatcog dattigeted cogen 125 <210> SEQ ID NO: 9	
125 <2115 SEQ 1D NO. 3 126 <2115 LENGTH: 25	
127 <212> TYPE: DNA	
128 <213> ORGANTSM: Unknown	
131 <223> OTHER INFORMATION: DELIVED TEST	
122 Z4005 SEQUENCE: 9	25
134 gggagatota gacatgataa galac	
137 <210> SEQ ID NO: 10	
138 <211> LENGTH: 27	
139 <212> TYPE: DNA	
140 <213> ORGANISM: Unknown	



RAW SEQUENCE LISTING DATE: 11/17/2603
PARENT APPLICATION: US/09/356,575D TIME: 09:26:45

142 <220> PEATURE: 143 <223> OTHER INFORMATION: Derived from Adenovirus 145 <400> SEQUENC: 10 146 aggagatoty tattgaaaty tytyggc 149 <210> SEO ID NO: 11 150 <211> LENGTH: 24	27
151 <212> FYPE: ONA 152 <213> ORGANISH: Unknown 154 <220> FEATURE: 155 <223> OTHER INFORMATION: Derived from Adenovirus 157 <400> SEQUENCE: 11 158 qqaqqetqca qtetecaacq qcqt 161 <210> SEQ 1D NO: 12 162 <211> LENGTH: 27 163 <212> TYPE: DNA	24
164 <213> ORGANISM: Unknown 166 <220> FEATURE: 167 <223> OTHER INFORMATION: Derived from Adenovirus 169 <400> SEQUENCE: 12 170 quqqqatcet caaateqtea etteeqt 173 <210> SEQ 1D NO: 13 174 <211> LENGTH: 27	27
175 <212> TYPE: DNA 176 <213> ORGANISM: Unknown 178 <220> PERTURE: 179 <223> OTHER INFORMATION: Derived from Adenovirus 181 <400> SEQUENCF: 13 182 gggqtctaga catcatcaat aatatac 185 <210> SEO ID NO: 14 186 <211> LENGTH: 32	27
187 <21.2> TYPE: DNA 188 <213> ORGANISM: Unknown 190 <220> FEATURE: 191 <223> OTHER INFORMATION: Derived from Adenovirus 193 <400> SEQUENCE: 14 194 ggcgaattcg tcgacatcat caataatata cc 197 <210> SEQ ID NO: 15 198 <211> LENGTH: 32	32
199 <212> TYPE: DNA 200 <213> ORGANISM: Unknown 202 <220> FEATURE: 203 <223> OTHER INFORMATION: Derived from Adenovirus 205 <400> SEQUENCE: 15 206 ggcgaattcg gtaccatcat caataatata cc 209 <210> SEO ID NO: 16 210 <211> LENGTH: 17	32
211 <212> TYPE: DNA 212 <213> ORGANISM: Unknown 214 <220> FEATURE:	

RAW SEQUENCE LISTING DATE: 11/17/26UC TIME: 09:26:43

Output Set. W. Care (	
217 <4005 SEMBERGE. 12 218 ctgtgtacac cgqcgca 221 <2105 SEM ID NO: 17 222 <2115 LENGTH: 50	17
223 <212> FYPE: DNA 221 <213> ORGANISM: Unknown 226 <220> FEATURE: 227 <223> OTHER INFORMATION: Derived from Adenovirus 229 <400> SEQUENCE: 17 230 ctacactgac ctaytgccgc ccgqgcaaag cccqggcgqc actaggtcag 233 <210> SEQ 1D NO: 18 234 <211> LENGTH: 50	50
235 <212> TYPE: DNA 236 <213> ORGANISM: Unknown 238 <220> FEATURE: 239 <223> OTHER INFORMATION: Derived from Adenovirus 241 <400> SEQUENCE: 18 242 qtacctqacc tagtgccgcc cgggctttyc ccgggcggca ctaggtcagt 245 <210> SEQ ID NO: 19 246 <211> LENGTH: 55	50
247 <212> TYPE: DNA 248 <213> ORGANISM: Unknown 250 <220> FEATURE: 251 <223> OTHER INFORMATION: Derived from Adenovirus 253 <400> SEQUENCE: 19 254 qtacattgac ctagtgcegc ccgggcaaag cccgggcggc actaggtcaa tcgat 257 <210> SEQ ID NO: 20 258 <211> LENGTH: 55	55
259 <212> TYPE: DNA 260 <21.3> ORGANISM: Unknown 262 <220> FEATURE: 263 <2223> OTHER INFORMATION: Derived from Adenovirus 265 <400> SEQUENCE: 20 266 gtacatcgat tgacctagtg ccgcccgggc tttgcccggg cggcactagg tcaat 269 <210> SEQ ID NO: 21 270 <211> LENGTH: 4900 271 <212> TYPE: DNA	55
271 <213> ORGANISM: Unknown 272 <220> FEATURE: 273 <223> OTHER INFORMATION: Derived from Adenovirus 274 <400> SEQUENCE: 21 278 ctttectact tagcastaca tetacestatt agteateset eggastate attaceatag tgatsgesst taggastate eaactigges gtogatages gtttgatte eaactigges gtesateset eaactigges gtttgattt ggatsgesst tetacaaaats eaacggast tetacaaaats eaacggast tecsaaaats eaacggast eagassecate eagass	60 120 180 240 300 360 420

RAW SEQUENCE LISTING PAILINI APPLICATION: US/09/356,575D

DAIL: 11/17/2000 FIME: 03:26:43

	180
292 gatcooggat cogglaging igcumutema agametgete etcaqiqqat gitycettia 4	540
292 quirecqqat ecqqiqqiqq iqeaaltaa uquategeri (acaitect gaatqleger egcaqiquea 294 ettelagiat cuagettgaa itectityig tideaitett gaatqleger egcaqiquea 294 ettelagiat cuagettgaa itecatique agargeeaad aacataaaqa aaqqeeegqe	500
294 cttctagtat caagettgaa iteettigig tidaattega aacataaaqa aaqqeeegqe 8 296 ttageattee qgtactqttg qtaaaatgga agaegeeaaa aacataaaqa aaqqeeegqe t	560
296 tragratice ggtactqttg qtaaaatqqa agatqttaataataaq ctatqaagaa 193 qecattctat cetetagagg argqaarege tggagagcaa etgcataaqag tgaucatcae 193 qecattctat cetetagagg argqaarege tagagatuca catategagg tgaucatcae	720
293 gecattetat cetetagagg atggaarege tygagageat estategagg tgaacateac 300 atacgecetg qtteetggaa eaattgettt tacagatgca catategagg tgaacateac	780
300 atacgecont qttcctqqaa caartgetti tacaquage detatgaaac gatatgaget 302 gtacgegqaa tacttcgaaa tgtccqtteg gttggcagaa getatgaaac gatatgaget	840
302 glacgoggaa tacttogaaa tgtooqtidy gloogaaactot ottoaattot ttatqooqqt 304 qaatacaaat cacagaatog togtatycag tgaaaactot ottoaattot ttatqooqqt	900
404 qaatacaaat cacagaatcy togtatycay tyaatacca aacqacattt ataatqaacq 306 gttgggcycg ffatttatcy qaqttqcayt tgcgcccgcg aacqacattt ataatqaacq	960
306 gttgggegeg ftatttateg gagttgeadt tytggetetgeg dattggtt ecaaaaaggg 308 tguattgete uacagtatga acatteegea geetacegta gtgtttgttt ecaaaaaggg 1	020
308 tquattgoto uacagtatga acarttegod geotactgu gegada tlattatoat 1 310 gttgoadadu uttttgaacg tgoadadaaa attuccaata atcoagdada tlattatoat 1	080
310 gitgcaaaaa attitgaacg igcaaaaada attaccaaca acgilogica catcleatet 1 312 ggattelaaa acggattaca agggattica gitgaatgtac acgilogica catcleatet 1	140
312 qgattelaaa acqqattacc agggatttaa tetqatetta acaaaacaat 1 314 aceteeggt tttaatqaat acqatttqt aceagaqtee tttqateqtq acaaaacaat 1	200
314 accteceggt tithatquat acquittiqt actagatete totagatete earteegea 1 316 tocactguta atquatteet etggatetae tgggttaect aagggtgtgg ecetteegea 1 316 tocactguta atquatteet etggatetee tgggatetet attitiggea atcamateut 1	260
316 tigractigata atgaintect ctigatetae tiggquetaet atgittitiggea atgainteat 1 318 tagaintige tigrigate tetrografie cagainteat atfittigga atgainteat 1	320
318 tagaactgcc tgcgtcagat tctcgcatgc catagatite untiltigaa tgtttactac 1 320 tccggatact gcgattttaa gtgttgttcc attccatcac ggttttggaa tgtttactac 1	380
320 tecaggatact gegattitaa gigtigitem alteedata systematig aagaagaget 1 322 aeteggatat tigatatgig gattiegagi egettaatg tatagattig aagaagaget 1	440
322 acteggatat ttgatatgtg gattecgagt cytetenary edgestagtac caaccetatt 1 324 gtttttaega teeetteagg attacaaaat teaaagtgeg ttgetagtac caaccetatt 1	L500
324 gtttttaega teeetteaga attacadaat teaaagagaga teatetaatt tacaegaaat 1 326 tteattette geeaaaagea etelgatiga caaataegat ttatetaatt tacaegaaat 1	1560
326 ttcattctte gecaaaagea etetgatta caattagat tetetaa aaegetteea 1 328 tgettetygg ggegeacete tttegaaaga agteggggaa geygttgeaa aaegetteea 1	L620
328 tgrttetygg ggegraeete titegaaaga aqtegygyaa geganaaga tietgattae 1 330 tetteeaggg atacgacaag qatatggget caetgagaet acateageta tietgaagegaa 1	1680
330 tottocaggg atacgacaag qatatgggot cactgagact ditocattit tigaagegaa 1332 accegagggg gatgataaac egggegeggt eggtaaagtt gitceattit tigaagegaa 1	1740
332 accognegge gatgatanac congregory togethant cananagyor antiatyty 1 334 ggttgtggat cuggatacog ggaanacgot gggcyttaat cananagyorg antiatytyt 1	1800
	1860
336 cagaggacct atgattatgt ccggttangt uddeduced tyggacgaag acgaacactt. 338 tgacaaggat ggatggctac attotggaga catagettac tyggacgaag acgaacactt.	1920
338 tgacaaggat ggatggctac attetygaga catagoras ggatatcagg tggcccccqc 340 cttcatagtt gaccgcttga agtctttaat taaatacaaa ggatatcagg tggcaggtct	1980
340 otteatagtt gaccgottga agtottaat Laantacudu gaccgoggog tggcaggtot 342 tgaattggaa togatattgt tacaacacco caacatotto gaccgoggog tggcaggtot gaccaggaagac	2040
342 tyaattiggaa tegatattigt tacaacacee cacattette gasgae acquaaagae 344 teecgaegat gacqeeqqia aactteecge egeegttigti gtrittiggage acquaaaagae 344 teecgaegat gacqeeqqia acquaattigga gtaacaacee egaaaaagtt	2100
344 tecogacyat gacgocygig aactteedyd cyceyteyte y gaacaaccy cyaaaaagtt 346 gatgacggaa aaagagateg tygattacgt cyceagteaa gtaacaaccy cyaaaaagtt 346 gatgacggaa aactegacyc	2160
	2220
348 gegeggagga gitgigitti tiggaegaayi deesdaagge ggaaagteea aattgiaaaa 350 aagaaaate agagagatee teataaagge caagaagge ggaaagteea aattgiaaaa 350 aagaaaate agagagatee teataaagge agagagtee aattgi aatgggggat ceccaactig	2280
352 tgtaactgta ttcagcgatg acgaaatte tagcaattagga tcagaaattt cacaaataaa	2340
352 tylaactyta ticagogaty acquaattot tayotattys kursylyst 354 titattycag ottataatgy tiaccaaataa agcaatagca toaccaaatti cacaaataaa 356 goattiitti cacigoatto tayttytyy tiytocaaac toatcaatyi alottatoat 356 goattiitti cacigoatto taytagaaga gotogaatto ylaatcatyy toatagotyt	2400
356 geathtittt cartgeatte tagttgtggt tightcaaat dataatg teatagetgt	2460
358 gtotggateg gategatere egggtatega strongerea catacgagee ggaageataa	2520
360 ttoctgtgtg aaattgttat cegettaataa getraetgas attaattgeg ttgegeteac	2580
362 agtgtaaagc ctggggtgcc taatgagtga reasgataga traatgaatc ggccaacgcg	2640
364 tgcccgcttt ccagtcggga adcttgtegt getteggette ctcgctcact gactcgctgc	2700
366 cggggagayg cggtttgcgt attgggggtatt cagetcactc aaaggcggta atacggttat	2760
368 geteggtegt teggetgegg tigageggter acategage aaaaggeeag caaaaggeea	2820
370 ccacaqaatc aqqqqataac ytayyaaagt tittocataq gctcgcccc cctgacqagc	2880
372 ggaaccgtaa aaaggccgcg ttgcttgcqt tgggaaaccc yacaggacta taaagatacc	2940
374 atcacaaaaa togacgotca agutagaga yayaya yaratacatat togaccetq cogottaccq	3000
376 aggegitted cooligaage leectegage regisgeget theteatage teacgetgia	3060
378 gatacetgic egectitele ettergaget geargetggg etgigtgee gaaceceeg	3120
380 ggtateteag tteggtgtag gtegttegte bearing to transfer tgagtecaae eeggtaagae	3180
382 ttcagcccga ccgctgcgcc ttatcccgcta dtaacaggat tagcagagcg aggtatgtag	3240
384 acquettate gecactigge geogretary stranguage chacactaga aggacagtat	3300
386 geggtgetae agagiteitg aagtggtigge claacingg edagttiggt agcietigat 388 tigglaietg egetetyetg aageeagita celteggaaa aagagitiggi agcietigat	3360
388 tigglatoty egetetyciy daycedyced booksys	

VERIFICATION SUMMARYDATE: 11/17/2500PATENT APPLICATION: US/09/356,575DTime: 09:26:44